RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/531,662_
Source: _PCT/10_
Date Processed by STIC: _4/26/05_

ENTERED

RAW SEQUENCE LISTING   DATE: 04/26/2005
PATENT APPLICATION: US/10/531,662   TIME: 15:44:30

Input Set : **A:\PCTGB2003004492 sequence listing.txt**
Output Set: **N:\CRF4\04262005\J531662.raw**

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      3 <110> APPLICANT: ISIS INNOVATION LIMITED
      5 <120> TITLE OF INVENTION: HYDROXYLASES AND MODULATORS THEREOF
      7 <130> FILE REFERENCE: 06843.0091
C-->  9 <140> CURRENT APPLICATION NUMBER: US/10/531,662
C--> 10 <141> CURRENT FILING DATE: 2005-04-15
     12 <150> PRIOR APPLICATION NUMBER: GB 0224102.4
     13 <151> PRIOR FILING DATE: 2002-10-16
     15 <150> PRIOR APPLICATION NUMBER: GB 0226598.1
     16 <151> PRIOR FILING DATE: 2002-11-14
     18 <160> NUMBER OF SEQ ID NOS: 32
     20 <170> SOFTWARE: PatentIn version 3.2
     22 <210> SEQ ID NO: 1
     23 <211> LENGTH: 14
     24 <212> TYPE: PRT
     25 <213> ORGANISM: Artificial sequence
     27 <220> FEATURE:
     28 <221> NAME/KEY: source
     29 <223> OTHER INFORMATION: Artificial peptide
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     34 1               5                  10
     37 <210> SEQ ID NO: 2
     38 <211> LENGTH: 8
     39 <212> TYPE: PRT
     40 <213> ORGANISM: Artificial sequence
     42 <220> FEATURE:
     43 <221> NAME/KEY: source
     44 <223> OTHER INFORMATION: Artificial peptide
     46 <400> SEQUENCE: 2
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     49 1               5
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     53 <211> LENGTH: 17
     54 <212> TYPE: PRT
     55 <213> ORGANISM: Artificial sequence
     57 <220> FEATURE:
     58 <221> NAME/KEY: source
     59 <223> OTHER INFORMATION: Artificial peptide
     62 <220> FEATURE:
     63 <221> NAME/KEY: misc_feature
     64 <222> LOCATION: (16)..(16)
     65 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except
asparagine
     67 <400> SEQUENCE: 3
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     73 Ala
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     78 <211> LENGTH: 58
     79 <212> TYPE: PRT
     80 <213> ORGANISM: Homo sapiens
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     84 Phe Asn Trp Asn Trp Ile Asn Lys Gln Gln Gly Lys Arg Gly Trp Gly
     85 1               5                   10                  15
     88 Gln Leu Thr Ser Asn Leu Leu Leu Ile Gly Met Glu Gly Asn Val Thr
     89               20                  25                  30
     92 Pro Ala His Tyr Asp Glu Gln Gln Asn Phe Phe Ala Gln Ile Lys Gly
     93           35                  40                  45
     96 Tyr Lys Arg Cys Ile Leu Phe Pro Pro Asp
     97     50                  55
    100 <210> SEQ ID NO: 5
    101 <211> LENGTH: 16
    102 <212> TYPE: PRT
    103 <213> ORGANISM: Drosophila melanogaster
    105 <400> SEQUENCE: 5
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    108 1               5                   10                  15
    111 <210> SEQ ID NO: 6
    112 <211> LENGTH: 42
    113 <212> TYPE: PRT
    114 <213> ORGANISM: Drosophila melanogaster
    116 <400> SEQUENCE: 6
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    119 1               5                   10                  15
    122 Ser Met His Lys Asp Pro Tyr Glu Asn Val Tyr Cys Val Ile Ser Gly
    123               20                  25                  30
    126 His Lys Asp Phe Val Leu Ile Pro Pro His
    127           35                  40
    130 <210> SEQ ID NO: 7
    131 <211> LENGTH: 14
    132 <212> TYPE: PRT
    133 <213> ORGANISM: Drosophila melanogaster
    135 <400> SEQUENCE: 7
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    138 1               5                   10
    141 <210> SEQ ID NO: 8
    142 <211> LENGTH: 43
    143 <212> TYPE: PRT
    144 <213> ORGANISM: Drosophila melanogaster
    146 <400> SEQUENCE: 8
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    149 1               5                   10                  15
    152 Ser Pro Met His Tyr Asp Pro Lys His Asn Leu Leu Cys Gln Val Phe
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157         35                40
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163 <213> ORGANISM: Homo sapiens
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168 1               5                10                15
171 Cys
175 <210> SEQ ID NO: 10
176 <211> LENGTH: 44
177 <212> TYPE: PRT
178 <213> ORGANISM: Homo sapiens
180 <400> SEQUENCE: 10
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183 1               5                10                15
186 Asp Phe His Ile Asp Phe Gly Gly Thr Ser Val Trp Tyr His Val Leu
187         20                25                30
190 Lys Gly Glu Lys Ile Phe Tyr Leu Ile Arg Pro Thr
191         35                40
194 <210> SEQ ID NO: 11
195 <211> LENGTH: 16
196 <212> TYPE: PRT
197 <213> ORGANISM: Caenorhabditis elegans
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202 1               5                10                15
205 <210> SEQ ID NO: 12
206 <211> LENGTH: 44
207 <212> TYPE: PRT
208 <213> ORGANISM: Caenorhabditis elegans
210 <400> SEQUENCE: 12
212 Pro Lys Val Glu Gln Phe Cys Leu Ala Gly Met Ala Gly Ser Tyr Thr
213 1               5                10                15
216 Asp Phe His Val Asp Phe Gly Gly Ser Ser Val Tyr Tyr His Ile Leu
217         20                25                30
220 Lys Gly Glu Lys Ile Phe Tyr Ile Ala Ala Pro Thr
221         35                40
224 <210> SEQ ID NO: 13
225 <211> LENGTH: 16
226 <212> TYPE: PRT
227 <213> ORGANISM: Caenorhabditis elegans
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236 <211> LENGTH: 35
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238 <213> ORGANISM: Caenorhabditis elegans
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247             20                25                30
250 Lys Gly Glu
251          35
254 <210> SEQ ID NO: 15
255 <211> LENGTH: 9
256 <212> TYPE: PRT
257 <213> ORGANISM: Caenorhabditis elegans
259 <400> SEQUENCE: 15
261 Lys Ile Phe Tyr Ile Ala Ala Pro Thr
262 1               5
265 <210> SEQ ID NO: 16
266 <211> LENGTH: 16
267 <212> TYPE: PRT
268 <213> ORGANISM: Drosophila melanogaster
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272 Glu Ile Val Arg Gln Ile Asp Trp Val Asp Val Val Trp Pro Lys Gln
273 1               5                10                15
276 <210> SEQ ID NO: 17
277 <211> LENGTH: 35
278 <212> TYPE: PRT
279 <213> ORGANISM: Drosophila melanogaster
281 <400> SEQUENCE: 17
283 Pro Lys Val Gln Lys Tyr Cys Leu Met Ser Val Lys Asn Cys Tyr Thr
284 1               5                10                15
287 Asp Phe His Ile Asp Phe Gly Gly Thr Ser Val Trp Tyr His Ile Leu
288             20                25                30
291 Arg Gly Ser
292          35
295 <210> SEQ ID NO: 18
296 <211> LENGTH: 9
297 <212> TYPE: PRT
298 <213> ORGANISM: Drosophila melanogaster
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302 Lys Val Phe Trp Leu Ile Pro Pro Thr
303 1               5
306 <210> SEQ ID NO: 19
307 <211> LENGTH: 18
308 <212> TYPE: PRT
309 <213> ORGANISM: Saccharomyces cerevisiae
311 <400> SEQUENCE: 19
313 Gln Asn Asp Leu Val Asp Lys Ile Trp Ser Phe Asn Gly His Leu Glu
314 1               5                10                15
317 Lys Val
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322 <211> LENGTH: 44
323 <212> TYPE: PRT
324 <213> ORGANISM: Saccharomyces cerevisiae
326 <400> SEQUENCE: 20
328 Pro Lys Val Thr Lys Tyr Ile Leu Met Ser Val Lys Asp Ala Tyr Thr
329 1               5                  10                  15
332 Asp Phe His Leu Asp Phe Ala Gly Thr Ser Val Tyr Tyr Asn Val Ile
333             20                  25                  30
336 Ser Gly Gln Lys Lys Phe Leu Leu Phe Pro Pro Thr
337         35                  40
340 <210> SEQ ID NO: 21
341 <211> LENGTH: 61
342 <212> TYPE: PRT
343 <213> ORGANISM: Rattus norvegicus
345 <400> SEQUENCE: 21
347 Lys Thr Asp Val Phe Gln Glu Val Met Trp Ser Asp Phe Gly Phe Pro
348 1               5                  10                  15
351 Gly Arg Asn Gly Gln Glu Ser Thr Leu Trp Ile Gly Ser Leu Gly Ala
352             20                  25                  30
355 His Thr Pro Cys His Leu Asp Ser Tyr Gly Cys Asn Leu Val Phe Gln
356             35                  40                  45
359 Val Gln Gly Arg Lys Arg Trp His Leu Phe Pro Pro Glu
360         50                  55                  60
363 <210> SEQ ID NO: 22
364 <211> LENGTH: 57
365 <212> TYPE: PRT
366 <213> ORGANISM: Caenorhabditis elegans
368 <400> SEQUENCE: 22
370 Phe Glu Asp Asp Leu Phe His Tyr Ala Asp Asp Lys Lys Arg Pro Pro
371 1               5                  10                  15
374 His Arg Trp Phe Val Met Gly Pro Ala Arg Ser Gly Thr Ala Ile His
375             20                  25                  30
378 Ile Asp Pro Leu Gly Thr Ser Ala Trp Asn Ser Leu Leu Gln Gly His
379             35                  40                  45
382 Lys Arg Trp Val Leu Ile Pro Pro Ile
383         50                  55
386 <210> SEQ ID NO: 23
387 <211> LENGTH: 60
388 <212> TYPE: PRT
389 <213> ORGANISM: Drosophila melanogaster
391 <400> SEQUENCE: 23
393 Thr Ile Leu Asp Tyr Val Asn Lys Asp Tyr Asn Ile Gln Ile Asp Gly
394 1               5                  10                  15
397 Val Asn Thr Ala Tyr Leu Tyr Phe Gly Met Trp Lys Thr Thr Phe Ala
398             20                  25                  30
401 Trp His Thr Glu Asp Met Asp Leu Tyr Ser Ile Asn Tyr Leu His Phe
402             35                  40                  45
405 Gly Ala Pro Lys Thr Trp Tyr Val Val Pro Pro Glu
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 04/26/2005
PATENT APPLICATION: US/10/531,662   TIME: 15:44:31

Input Set,: **A:\PCTGB2003004492 sequence listing.txt**
Output Set: **N:\CRF4\04262005\J531662.raw**

<u>Please Note:</u>
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 16

VERIFICATION SUMMARY   DATE: 04/26/2005
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Input Set : **A:\PCTGB2003004492 sequence listing.txt**
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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:69 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0